## SEQUENCE LISTING

<110>	Oft, Mar McClanah		ill K.		•								
<120>	USES OF	IL-23 AG	ONISTS A	ND ANT	'AGONIST	S; REL	ATED F	REAGE	ENTS				
<130>	DX06022U	S01											
<150> <151>	U.S. 60/453,672 2003-03-10												
<160>	11												
<170>	PatentIn	version	3.2										
<210><211><211><212><213>	DNA	iens											
<220> <221> <222>	CDS (1)(56	7)											
<220> <221> <222>													
<400> atg ct Met Le -2	1 g ggg agc u Gly Ser 0	aga gct Arg Ala	gta atg Val Met -15	ctg c Leu L	tg ttg eu Leu	ctg ct Leu Le	g ccc u Pro	tgg Trp	aca Thr	48			
gct ca Ala Gl -5	g ggc aga n Gly Arg	gct gtg Ala Val -1 1	cct ggg Pro Gly	ggc a Gly S	er Ser	cct go Pro Al	c tgg a Trp	act Thr 10	cag Gln	96			
	g cag ctt n Gln Leu 15									144			
cca ct Pro Le	a gtg gga u Val Gly 30	cac atg His Met	gat cta Asp Leu 35	aga g Arg G	gaa gag Slu Glu	gga ga Gly As 40	p Glu	gag Glu	act Thr	192			
	t gat gtt n Asp Val									240			
	c agg gac u Arg Asp									288			
	t ttt tat e Phe Tyr		_	Gly S						336			
	t ctg ctc r Leu Leu 95									384			

	ggc Gly															433	2
_	cag Gln 125			_		_		_	_			_	_			480	0
	cgc Arg															528	8
	cgg Arg												taa			576	0
<21 <21 <21 <21	1> 1 2> 1		sapi	iens											٠		
<40	0 > 2	2															
Met	Leu -20	Gly	Ser	Arg	Ala	Val -15	Met	Leu	Leu	Leu	Leu -10	Leu	Pro	Trp	Thr		
Ala -5	Gln	Gly	Arg	Ala -1	Val 1	Pro	Gly	Gly	Ser 5	Ser	Pro	Ala	Trp	Thr 10	Gln		
Cys	Gln	Gln	Leu 15	Ser	Gln	Lys	Leu	Cys 20	Thr	Leu	Ala	Trp	Ser 25	Ala	His		
Pro	Leu	Val 30	Gly	His	Met	Asp	Leu 35	Arg	Glu	Glu	Gly	Asp 40	Glu	Glu	Thr		
Thr	Asn 45	Asp	Val	Pro	His	Ile 50	Gln	Cys	Gly	Asp	Gly 55	Cys	Asp	Pro	Gln		
Gly 60	Leu	Arg	Asp	Asn	Ser 65	Gln	Phe	Cys	Leu	Gln 70	Arg	Ile	His	Gln	Gly 75		
Leu	Ile	Phe	Tyr	Glu 80	Lys	Leu	Leu	Gly	Ser 85	Asp	Ile	Phe	Thr	Gly 90	Glu		
Pro	Ser	Leu	Leu 95	Pro	Asp	Ser	Pro	Val 100	Ala	Gln	Leu	His	Ala 105	Ser	Leu		
Leu	Gly	Leu 110	Ser	Gln	Leu	Leu	Gln 115	Pro	Glu	Gly	His	His 120	Trp	Glu	Thr		

Gln Gln Ile Pro Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu

125 130 135

Leu Arg Phe Lys Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala Ala Arg Val Phe Ala His Gly Ala Ala Thr Leu Ser Pro <210> 3 1203 <211> <212> DNA <213> Mus musculus <220> <221> CDS <222> (113)..(700) <220> <221> mat\_peptide <222> (176)..(700) <400> 3 cqcttagaag tcggactaca gagttagact cagaaccaaa ggaggtggat agggggtcca caggcctggt gcagatcaca gagccagcca gatctgagaa gcagggaaca ag atg ctg 118 Met Leu gat tgc aga gca gta ata atg cta tgg ctg ttg ccc tgg gtc act cag 166 Asp Cys Arg Ala Val Ile Met Leu Trp Leu Leu Pro Trp Val Thr Gln -15 -10 gge etg get gtg eet agg agt age agt eet gae tgg get eag tge eag 214 Gly Leu Ala Val Pro Arg Ser Ser Ser Pro Asp Trp Ala Gln Cys Gln cag etc tet egg aat etc tge atg eta gee tgg aac gea eat gea eea 262 Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His Ala Pro 20 310 qcq qqa cat atg aat cta cta aga gaa gaa gag gat gaa gag act aaa Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu Thr Lys 35 358 aat aat gtg ccc cgt atc cag tgt gaa gat ggt tgt gac cca caa gga Asn Asn Val Pro Arq Ile Gln Cys Glu Asp Gly Cys Asp Pro Gln Gly 50 55 ctc aag gac aac agc cag ttc tgc ttg caa agg atc cgc caa ggt ctg 406 Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln Gly Leu 65 454 gct ttt tat aag cac ctg ctt gac tct gac atc ttc aaa ggg gag cct Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly Glu Pro 80 85 90

gct cta ctc cct gat agc ccc atg gag caa ctt cac acc tcc cta cta

Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser Leu Leu

502

	95					100					105					
														acc Thr		550
_	_		_	_	_		_	_	_		_	_		ctt Leu 140		598
														gct Ala		646
														gtg Val		694
Thr	gct Ala 175	taaq	ggat	gcc (	caggt	tcc	ca to	ggcta	accat	gat	aaga	acta	atc	tatca	agc	750
cca	gacat	ct a	acca	gttaa	at ta	acco	catta	a gga	actt	gtgc	tgtt	ctt	gtt	tcgtt	tgttt	810
tgc	gtgaa	agg g	gcaa	ggac	ac ca	attat	taaa	a gag	gaaaa	agaa	acaa	acco	cca 🤅	gagca	aggcag	870
ctg	gctag	gag a	aaag	gagct	g ga	agaag	gaaga	a ata	aaagt	ctc	gago	ccctt	gg (	cctt	ggaagc	930
999	caago	cag (	ctgc	gtgg	cc to	gagg	ggaag	g gg	ggcgg	gtgg	cato	gaga	aaa	ctgt	gagaaa	990
acc	caga	gca 1	tcag	aaaa	ag to	gagco	ccago	g cti	ttgg	ccat	tato	tgta	aag	aaaa	acaaga	1050
aaa	3999	aac a	atta	tact	t c	tgg	gtggd	tca	aggga	aaat	gtg	cagat	gc a	acagt	cactcc	1110
aga	cagca	agc 1	tctg	tacci	g co	ctgct	ctgt	ccc	ctca	gttc	taad	cagaa	atc	tagto	cactaa	1170
gaa	ctaa	cag g	gact	accaa	at a	cgaad	ctgad	c aaa	a							1203

<210> 4 <211> 196 <212> PRT <213> Mus musculus

<400> 4

Met Leu Asp Cys Arg Ala Val Ile Met Leu Trp Leu Leu Pro Trp Val -20 -15

Thr Gln Gly Leu Ala Val Pro Arg Ser Ser Pro Asp Trp Ala Gln -1 1 -5

Cys Gln Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His 20

Ala Pro Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu 35 40

Thr Lys Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro 45 Gln Gly Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln Gly Leu Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly Glu Pro Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser 100 Leu Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu 115 Thr Gln Gln Met Pro Ser Leu Ser Ser Gln Gln Trp Gln Arg Pro 130 Leu Leu Arg Ser Lys Ile Leu Arg Ser Leu Gln Ala Phe Leu Ala Ile 150 Ala Ala Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu Val Pro Thr Ala 175 <210> 5 <211> 2859 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (119)..(2005) <220> mat\_peptide <221> (188)..(2005) <400> 5 gtggtacggg aattccattg tgttgggcag ccaacaaggg tggcagcctg gctctgaagt 60 ggaattatgt gcttcaaaca ggttgaaaga gggaaacagt cttttcctgc ttccagac 118 atg aat cak gtc act att caa tgg gat gca gta ata gcc ctt tac ata 166 Met Asn Xaa Val Thr Ile Gln Trp Asp Ala Val Ile Ala Leu Tyr Ile -15 ctc ttc agc tgg tgt cat gga gga att aca aat ata aac tgc tct ggc 214 Leu Phe Ser Trp Cys His Gly Gly Ile Thr Asn Ile Asn Cys Ser Gly
-5 -1 1 5

cac His 10	atc Ile	tgg Trp	gta Val	gaa Glu	cca Pro 15	gcc Ala	aca Thr	att Ile	ttt Phe	aag Lys 20	atg Met	ggt Gly	atg Met	aat Asn	atc Ile 25	262
														aaa Lys 40		310
cat His	ttt Phe	tat Tyr	aaa Lys 45	aat Asn	ggc Gly	atc Ile	aaa Lys	gaa Glu 50	aga Arg	ttt Phe	caa Gln	atc Ile	aca Thr 55	agg Arg	att Ile	358
aat Asn	aaa Lys	aca Thr 60	aca Thr	gct Ala	cgg Arg	ctt Leu	tgg Trp 65	tat Tyr	aaa Lys	aac Asn	ttt Phe	ctg Leu 70	gaa Glu	cca Pro	cat His	406
gct Ala	tct Ser 75	atg Met	tac Tyr	tgc Cys	act Thr	gct Ala 80	gaa Glu	tgt Cys	ccc Pro	aaa Lys	cat His 85	ttt Phe	caa Gln	gag Glu	aca Thr	454
_		_			_						_		_	att Ile		502
														act Thr 120		550
acc Thr	tgg Trp	aat Asn	gct Ala 125	rgg Xaa	aag Lys	ctc Leu	acc Thr	tac Tyr 130	ata Ile	gac Asp	aca Thr	aaa Lys	tac Tyr 135	gtg Val	gta Val	598
														acc Thr		646
agc Ser	tat Tyr 155	att Ile	aac Asn	atc Ile	tcc Ser	act Thr 160	gat Asp	tca Ser	tta Leu	caa Gln	ggt Gly 165	ggc Gly	aag Lys	aag Lys	tac Tyr	694
														tca Ser		742
caa Gln	ctg Leu	caa Gln	att Ile	cac His 190	ctg Leu	gat Asp	gat Asp	ata Ile	gtg Val 195	ata Ile	cct Pro	tct Ser	gca Ala	gcc Ala 200	gtc Val	790
														ata Ile		838
tat Tyr	tgg Trp	gat Asp 220	agt Ser	caa Gln	aca Thr	aca Thr	att Ile 225	gaa Glu	aag Lys	gtt Val	tcc Ser	tgt Cys 230	gaa Glu	atg Met	aga Arg	886
tac Tyr	aag Lys 235	gct Ala	aca Thr	aca Thr	aac Asn	caa Gln 240	act Thr	tgg Trp	aat Asn	gtt Val	aaa Lys 245	gaa Glu	ttt Phe	gac Asp	acc Thr	934
aat	ttt	aca	tat	gtg	caa	cag	tca	gaa	ttc	tac	ttg	gag	cca	aac	att	982

Asn 250	Phe	Thr	Tyr	Val	Gln 255	Gln	Ser	Glu	Phe	Tyr 260	Leu	Glu	Pro	Asn	Ile 265	
aag Lys	tac Tyr	gta Val	ttt Phe	caa Gln 270	gtg Val	aga Arg	tgt Cys	caa Gln	gaa Glu 275	aca Thr	ggc Gly	aaa Lys	agg Arg	tac Tyr 280	tgg Trp	1030
cag Gln	cct Pro	tgg Trp	agt Ser 285	tca Ser	ccg Pro	ttt Phe	ttt Phe	cat His 290	aaa Lys	aca Thr	cct Pro	gaa Glu	aca Thr 295	gtt Val	ccc Pro	1078
cag Gln	gtc Val	aca Thr 300	tca Ser	aaa Lys	gca Ala	ttc Phe	caa Gln 305	cat His	gac Asp	aca Thr	tgg Trp	aat Asn 310	tct Ser	G1y 999	cta Leu	1126
aca Thr	gtt Val 315	gct Ala	tcc Ser	atc Ile	tct Ser	aca Thr 320	gly aaa	cac His	ctt Leu	act Thr	tct Ser 325	gac Asp	aac Asn	aga Arg	gga Gly	1174
gac Asp 330	att Ile	gga Gly	ctt Leu	tta Leu	ttg Leu 335	gga Gly	atg Met	atc Ile	gtc Val	ttt Phe 340	gct Ala	gtt Val	atg Met	ttg Leu	tca Ser 345	1222
att Ile	ctt Leu	tct Ser	ttg Leu	att Ile 350	ggg Gly	ata Ile	ttt Phe	aac Asn	aga Arg 355	tca Ser	ttc Phe	cga Arg	act Thr	360 Gly 393	att Ile	1270
aaa Lys	aga Arg	agg Arg	atc Ile 365	tta Leu	ttg Leu	tta Leu	ata Ile	cca Pro 370	aag Lys	tgg Trp	ctt Leu	tat Tyr	gaa Glu 375	gat Asp	att Ile	1318
cct Pro	aat Asn	atg Met 380	aaa Lys	aac Asn	agc Ser	aat Asn	gtt Val 385	gtg Val	aaa Lys	atg Met	cta Leu	cag Gln 390	gaa Glu	aat Asn	agt Ser	1366
gaa Glu	ctt Leu 395	atg Met	aat Asn	aat Asn	aat Asn	tcc Ser 400	agt Ser	gag Glu	cag Gln	gtc Val	cta Leu 405	tat Tyr	gtt Val	gat Asp	ccc Pro	1414
atg Met 410	att Ile	aca Thr	gag Glu	ata Ile	aaa Lys 415	gaa Glu	atc Ile	ttc Phe	atc Ile	cca Pro 420	gaa Glu	cac His	aag Lys	cct Pro	aca Thr 425	1462
gac Asp	tac Tyr	aag Lys	aag Lys	gag Glu 430	aat Asn	aca Thr	gga Gly	ccc Pro	ctg Leu 435	gag Glu	aca Thr	aga Arg	gac Asp	tac Tyr 440	ccg Pro	1510
caa Gln	aac Asn	tcg Ser	cta Leu 445	ttc Phe	gac Asp	aat Asn	act Thr	aca Thr 450	gtt Val	gta Val	tat Tyr	att Ile	cct Pro 455	gat Asp	ctc Leu	1558
aac Asn	act Thr	gga Gly 460	tat Tyr	aaa Lys	ccc Pro	caa Gln	att Ile 465	tca Ser	aat Asn	ttt Phe	ctg Leu	cct Pro 470	gag Glu	gga Gly	agc Ser	1606
cat His	ctc Leu 475	agc Ser	aat Asn	aat Asn	aat Asn	gaa Glu 480	att Ile	act Thr	tcc Ser	tta Leu	aca Thr 485	ctt Leu	aaa Lys	cca Pro	cca Pro	1654
gtt Val 490	gat Asp	tcc Ser	tta Leu	gac Asp	tca Ser 495	gga Gly	aat Asn	aat Asn	ccc Pro	agg Arg 500	tta Leu	caa Gln	aag Lys	cat His	cct Pro 505	1702

aat ttt gct ttt tct gtt tca agt gtg aat tca cta agc aac aca ata Asn Phe Ala Phe Ser Val Ser Ser Val Asn Ser Leu Ser Asn Thr Ile 510 515 520	1750
ttt ctt gga gaa tta agc ctc ata tta aat caa gga gaa tgc agt tct Phe Leu Gly Glu Leu Ser Leu Ile Leu Asn Gln Gly Glu Cys Ser Ser 525 . 530 535	1798
cct gac ata caa aac tca gta gag gag gaa acc acc atg ctt ttg gaa Pro Asp Ile Gln Asn Ser Val Glu Glu Glu Thr Thr Met Leu Leu Glu 540 545 550	1846
aat gat tca ccc agt gaa act att cca gaa cag acc ctg ctt cct gat Asn Asp Ser Pro Ser Glu Thr Ile Pro Glu Gln Thr Leu Leu Pro Asp 555 560 565	1894
gaa ttt gtc tcc tgt ttg ggg atc gtg aat gag gag ttg cca tct att Glu Phe Val Ser Cys Leu Gly Ile Val Asn Glu Glu Leu Pro Ser Ile 570 575 580 585	1942
aat act tat ttt cca caa aat att ttg gaa agc cac ttc aat agg att Asn Thr Tyr Phe Pro Gln Asn Ile Leu Glu Ser His Phe Asn Arg Ile 590 595 600	1990
tca ctc ttg gaa aag tagagctgtg tggtcaaaat caatatgaga aagctgcctt Ser Leu Leu Glu Lys 605	2045
gcaatctgaa cttgggtttt ccctgcaata gaaattgaat tctgcctctt tttgaaaaaa	2105
atgtattcac atacaaatct tcacatggac acatgttttc atttcccttg gataaatacc	2165
taggtagggg attgctgggc catatgataa gcatatgttt cagttctacc aatcttgttt	2225
ccagagtagt gacatttctg tgctcctacc atcaccatgt aagaattccc gggagctcca	2285
tgccttttta attttagcca ttcttctgcc tmatttctta aaattagaga attaaggtcc	2345
cgaaggtgga acatgcttca tggtcacaca tacaggcaca aaaacagcat tatgtggacg	2405
cctcatgtat tttttataga gtcaactatt tcctctttat tttccctcat tgaaagatgc	2465
aaaacagctc tctattgtgt acagaaaggg taaataatgc aaaatacctg gtagtaaaat	2525
aaatgctgaa aattttcctt taaaatagaa tcattaggcc aggcgtggtg gctcatgctt	2585
gtaatcccag cactttggta ggctgaggtr ggtggatcac ctgaggtcag gagttcgagt	2645
ccagcctggc caatatgctg aaaccctgtc tctactaaaa ttacaaaaat tagccggcca	2705
tggtggcagg tgcttgtaat cccagctact tgggaggctg aggcaggaga atcacttgaa	2765
ccaggaaggc agaggttgca ctgagctgag attgtgccac tgcactccag cctgggcaac	2825
aagagcaaaa ctctgtctgg aaaaaaaaaa aaaa	2859

<sup>&</sup>lt;210> 6 <211> 629 <212> PRT <213> Homo sapiens

<220> <221> misc feature <222> (-21)..(-21) The 'Xaa' at location -21 stands for Gln, or His. <220> misc\_feature <221> (126)...(126)<222> The 'Xaa' at location 126 stands for Gly, or Arg. <400> 6 Met Asn Xaa Val Thr Ile Gln Trp Asp Ala Val Ile Ala Leu Tyr Ile -15 Leu Phe Ser Trp Cys His Gly Gly Ile Thr Asn Ile Asn Cys Ser Gly His Ile Trp Val Glu Pro Ala Thr Ile Phe Lys Met Gly Met Asn Ile Ser Ile Tyr Cys Gln Ala Ala Ile Lys Asn Cys Gln Pro Arg Lys Leu His Phe Tyr Lys Asn Gly Ile Lys Glu Arg Phe Gln Ile Thr Arg Ile Asn Lys Thr Thr Ala Arg Leu Trp Tyr Lys Asn Phe Leu Glu Pro His Ala Ser Met Tyr Cys Thr Ala Glu Cys Pro Lys His Phe Gln Glu Thr Leu Ile Cys Gly Lys Asp Ile Ser Ser Gly Tyr Pro Pro Asp Ile Pro Asp Glu Val Thr Cys Val Ile Tyr Glu Tyr Ser Gly Asn Met Thr Cys Thr Trp Asn Ala Xaa Lys Leu Thr Tyr Ile Asp Thr Lys Tyr Val Val 130 135 His Val Lys Ser Leu Glu Thr Glu Glu Glu Gln Gln Tyr Leu Thr Ser 140 Ser Tyr Ile Asn Ile Ser Thr Asp Ser Leu Gln Gly Gly Lys Lys Tyr 155 160 165

Leu Val Trp Val Gln Ala Ala Asn Ala Leu Gly Met Glu Glu Ser Lys

170	175	180	185

Gln Leu Gln Ile His Leu Asp Asp Ile Val Ile Pro Ser Ala Ala Val 190 195 200

Ile Ser Arg Ala Glu Thr Ile Asn Ala Thr Val Pro Lys Thr Ile Ile 205 210 215

Tyr Trp Asp Ser Gln Thr Thr Ile Glu Lys Val Ser Cys Glu Met Arg 220 225 230

Tyr Lys Ala Thr Thr Asn Gln Thr Trp Asn Val Lys Glu Phe Asp Thr 235 240 245

Asn Phe Thr Tyr Val Gln Gln Ser Glu Phe Tyr Leu Glu Pro Asn Ile 250 255 260 265

Lys Tyr Val Phe Gln Val Arg Cys Gln Glu Thr Gly Lys Arg Tyr Trp 270 275 280

Gln Pro Trp Ser Ser Pro Phe Phe His Lys Thr Pro Glu Thr Val Pro 285 290 295

Gln Val Thr Ser Lys Ala Phe Gln His Asp Thr Trp Asn Ser Gly Leu 300 305 310

Thr Val Ala Ser Ile Ser Thr Gly His Leu Thr Ser Asp Asn Arg Gly 315 320 325

Asp Ile Gly Leu Leu Gly Met Ile Val Phe Ala Val Met Leu Ser 330 345

Ile Leu Ser Leu Ile Gly Ile Phe Asn Arg Ser Phe Arg Thr Gly Ile 350 355 360

Lys Arg Arg Ile Leu Leu Leu Ile Pro Lys Trp Leu Tyr Glu Asp Ile 365 370 375

Pro Asn Met Lys Asn Ser Asn Val Val Lys Met Leu Gln Glu Asn Ser 380 385 390

Glu Leu Met Asn Asn Ser Ser Glu Gln Val Leu Tyr Val Asp Pro 395 400.

Met Ile Thr Glu Ile Lys Glu Ile Phe Ile Pro Glu His Lys Pro Thr 410 415 420 425

Asp Tyr Lys Lys Glu Asn Thr Gly Pro Leu Glu Thr Arg Asp Tyr Pro 430 435 440

Gln Asn Ser Leu Phe Asp Asn Thr Thr Val Val Tyr Ile Pro Asp Leu 445 450 455

Asn Thr Gly Tyr Lys Pro Gln Ile Ser Asn Phe Leu Pro Glu Gly Ser 460 465 470

His Leu Ser Asn Asn Glu Ile Thr Ser Leu Thr Leu Lys Pro Pro 475 480 485

Val Asp Ser Leu Asp Ser Gly Asn Asn Pro Arg Leu Gln Lys His Pro 490 495 500 505

Asn Phe Ala Phe Ser Val Ser Ser Val Asn Ser Leu Ser Asn Thr Ile 510 515 520

Phe Leu Gly Glu Leu Ser Leu Ile Leu Asn Gln Gly Glu Cys Ser Ser 525 530 535

Pro Asp Ile Gln Asn Ser Val Glu Glu Glu Thr Thr Met Leu Leu Glu 540 545 550

Asn Asp Ser Pro Ser Glu Thr Ile Pro Glu Gln Thr Leu Leu Pro Asp 555 560 565

Glu Phe Val Ser Cys Leu Gly Ile Val Asn Glu Glu Leu Pro Ser Ile 570 575 580 585

Asn Thr Tyr Phe Pro Gln Asn Ile Leu Glu Ser His Phe Asn Arg Ile 590 595 600

Ser Leu Leu Glu Lys 605

<210> 7

<211> 862

<212> PRT

<213> Homo sapiens

<400> 7

Met Ala His Thr Phe Arg Gly Cys Ser Leu Ala Phe Met Phe Ile Ile 1 5 10 15

Thr Trp Leu Leu Ile Lys Ala Lys Ile Asp Ala Cys Lys Arg Gly Asp 20 25 30

Val Thr Val Lys Pro Ser His Val Ile Leu Leu Gly Ser Thr Val Asn 35 40 45

Ile Thr Cys Ser Leu Lys Pro Arg Gln Gly Cys Phe His Tyr Ser Arg 50 55 60

Arg Asn Lys Leu Ile Leu Tyr Lys Phe Asp Arg Arg Ile Asn Phe His 70 75 80

His Gly His Ser Leu Asn Ser Gln Val Thr Gly Leu Pro Leu Gly Thr 85 90 95

Thr Leu Phe Val Cys Lys Leu Ala Cys Ile Asn Ser Asp Glu Ile Gln
100 105 110

Ile Cys Gly Ala Glu Ile Phe Val Gly Val Ala Pro Glu Gln Pro Gln 115 120 125

Asn Leu Ser Cys Ile Gln Lys Gly Glu Gln Gly Thr Val Ala Cys Thr 130 135 140

Trp Glu Arg Gly Arg Asp Thr His Leu Tyr Thr Glu Tyr Thr Leu Gln 145 150 155 160

Leu Ser Gly Pro Lys Asn Leu Thr Trp Gln Lys Gln Cys Lys Asp Ile 165 170 175

Tyr Cys Asp Tyr Leu Asp Phe Gly Ile Asn Leu Thr Pro Glu Ser Pro 180 185 190

Glu Ser Asn Phe Thr Ala Lys Val Thr Ala Val Asn Ser Leu Gly Ser 195 200 205

Ser Ser Ser Leu Pro Ser Thr Phe Thr Phe Leu Asp Ile Val Arg Pro 210 215 220

Leu Pro Pro Trp Asp Ile Arg Ile Lys Phe Gln Lys Ala Ser Val Ser 225 230 235 240

Arg Cys Thr Leu Tyr Trp Arg Asp Glu Gly Leu Val Leu Leu Asn Arg

Leu Arg Tyr Arg Pro Ser Asn Ser Arg Leu Trp Asn Met Val Asn Val 260 265 270

Thr Lys Ala Lys Gly Arg His Asp Leu Leu Asp Leu Lys Pro Phe Thr 275 Glu Tyr Glu Phe Gln Ile Ser Ser Lys Leu His Leu Tyr Lys Gly Ser Trp Ser Asp Trp Ser Glu Ser Leu Arg Ala Gln Thr Pro Glu Glu Glu 310 315 Pro Thr Gly Met Leu Asp Val Trp Tyr Met Lys Arg His Ile Asp Tyr Ser Arg Gln Gln Ile Ser Leu Phe Trp Lys Asn Leu Ser Val Ser Glu 345 Ala Arg Gly Lys Ile Leu His Tyr Gln Val Thr Leu Gln Glu Leu Thr Gly Gly Lys Ala Met Thr Gln Asn Ile Thr Gly His Thr Ser Trp Thr 375 Thr Val Ile Pro Arg Thr Gly Asn Trp Ala Val Ala Val Ser Ala Ala Asn Ser Lys Gly Ser Ser Leu Pro Thr Arg Ile Asn Ile Met Asn Leu 405 410 Cys Glu Ala Gly Leu Leu Ala Pro Arg Gln Val Ser Ala Asn Ser Glu 420 Gly Met Asp Asn Ile Leu Val Thr Trp Gln Pro Pro Arg Lys Asp Pro 435 Ser Ala Val Gln Glu Tyr Val Val Glu Trp Arg Glu Leu His Pro Gly 450 Gly Asp Thr Gln Val Pro Leu Asn Trp Leu Arg Ser Arg Pro Tyr Asn 475 470 Val Ser Ala Leu Ile Ser Glu Asn Ile Lys Ser Tyr Ile Cys Tyr Glu 485 Ile Arg Val Tyr Ala Leu Ser Gly Asp Gln Gly Cys Ser Ser Ile Leu Gly Asn Ser Lys His Lys Ala Pro Leu Ser Gly Pro His Ile Asn

520

Ala Ile Thr Glu Glu Lys Gly Ser Ile Leu Ile Ser Trp Asn Ser Ile 530 535 Pro Val Gln Glu Gln Met Gly Cys Leu Leu His Tyr Arg Ile Tyr Trp 550 555 Lys Glu Arg Asp Ser Asn Ser Gln Pro Gln Leu Cys Glu Ile Pro Tyr Arg Val Ser Gln Asn Ser His Pro Ile Asn Ser Leu Gln Pro Arg Val Thr Tyr Val Leu Trp Met Thr Ala Leu Thr Ala Ala Gly Glu Ser Ser His Gly Asn Glu Arg Glu Phe Cys Leu Gln Gly Lys Ala Asn Trp Met Ala Phe Val Ala Pro Ser Ile Cys Ile Ala Ile Ile Met Val Gly Ile Phe Ser Thr His Tyr Phe Gln Gln Lys Val Phe Val Leu Leu Ala Ala Leu Arg Pro Gln Trp Cys Ser Arg Glu Ile Pro Asp Pro Ala Asn Ser 670 665 660 Thr Cys Ala Lys Lys Tyr Pro Ile Ala Glu Glu Lys Thr Gln Leu Pro 675 680 Leu Asp Arg Leu Leu Ile Asp Trp Pro Thr Pro Glu Asp Pro Glu Pro 690 695 Leu Val Ile Ser Glu Val Leu His Gln Val Thr Pro Val Phe Arg His Pro Pro Cys Ser Asn Trp Pro Gln Arg Glu Lys Gly Ile Gln Gly His 730 Gln Ala Ser Glu Lys Asp Met Met His Ser Ala Ser Ser Pro Pro

Pro Arg Ala Leu Gln Ala Glu Ser Arg Gln Leu Val Asp Leu Tyr Lys

Val Leu Glu Ser Arg Gly Ser Asp Pro Lys Pro Glu Asn Pro Ala Cys 770 775 780

Pro Trp Thr Val Leu Pro Ala Gly Asp Leu Pro Thr His Asp Gly Tyr 785 790 795 800

Leu Pro Ser Asn Ile Asp Asp Leu Pro Ser His Glu Ala Pro Leu Ala 805 810 815

Asp Ser Leu Glu Glu Leu Glu Pro Gln His Ile Ser Leu Ser Val Phe 820 825 830

Pro Ser Ser Ser Leu His Pro Leu Thr Phe Ser Cys Gly Asp Lys Leu 835 840 845

Thr Leu Asp Gln Leu Lys Met Arg Cys Asp Ser Leu Met Leu 850 855 860

<210> 8

<211> 328

<212> PRT

<213> Homo sapiens

<400> 8

Met Cys His Gln Gln Leu Val Ile Ser Trp Phe Ser Leu Val Phe Leu 1 5 10 15

Ala Ser Pro Leu Val Ala Ile Trp Glu Leu Lys Lys Asp Val Tyr Val 20 25 30

Val Glu Leu Asp Trp Tyr Pro Asp Ala Pro Gly Glu Met Val Val Leu 35 40 45

Thr Cys Asp Thr Pro Glu Glu Asp Gly Ile Thr Trp Thr Leu Asp Gln 50 55 60

Ser Ser Glu Val Leu Gly Ser Gly Lys Thr Leu Thr Ile Gln Val Lys 65 70 75 80

Glu Phe Gly Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly Glu Val 85 90 95

Leu Ser His Ser Leu Leu Leu His Lys Lys Glu Asp Gly Ile Trp
100 105 110

Ser Thr Asp Ile Leu Lys Asp Gln Lys Glu Pro Lys Asn Lys Thr Phe 115 120 125 Leu Arg Cys Glu Ala Lys Asn Tyr Ser Gly Arg Phe Thr Cys Trp Trp 130 135 140

Leu Thr Thr Ile Ser Thr Asp Leu Thr Phe Ser Val Lys Ser Ser Arg 145 150 155 160

Gly Ser Ser Asp Pro Gln Gly Val Thr Cys Gly Ala Ala Thr Leu Ser 165 170 175

Ala Glu Arg Val Arg Gly Asp Asn Lys Glu Tyr Glu Tyr Ser Val Glu 180 185 190

Cys Gln Glu Asp Ser Ala Cys Pro Ala Ala Glu Glu Ser Leu Pro Ile 195 200 205

Glu Val Met Val Asp Ala Val His Lys Leu Lys Tyr Glu Asn Tyr Thr 210 215 220

Ser Ser Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro Lys Asn 225 230 235 240

Leu Gln Leu Lys Pro Leu Lys Asn Ser Arg Gln Val Glu Val Ser Trp 245 250 255

Glu Tyr Pro Asp Thr Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Thr 260 265 270

Phe Cys Val Gln Val Gln Gly Lys Ser Lys Arg Glu Lys Lys Asp Arg 275 280 285

Val Phe Thr Asp Lys Thr Ser Ala Thr Val Ile Cys Arg Lys Asn Ala 290 295 300

Ser Ile Ser Val Arg Ala Gln Asp Arg Tyr Tyr Ser Ser Ser Trp Ser 305 310 315 320

Glu Trp Ala Ser Val Pro Cys Ser 325

<210> 9

<211> 335

<212> PRT

<213> Mus musculus

<400> 9

Met Cys Pro Gln Lys Leu Thr Ile Ser Trp Phe Ala Ile Val Leu Leu 1 5 10 15

Val Ser Pro Leu Met Ala Met Trp Glu Leu Glu Lys Asp Val Tyr Val 20 25 30

Val Glu Val Asp Trp Thr Pro Asp Ala Pro Gly Glu Thr Val Asn Leu  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Thr Cys Asp Thr Pro Glu Glu Asp Asp Ile Thr Trp Thr Ser Asp Gln 50 55 60

Arg His Gly Val Ile Gly Ser Gly Lys Thr Leu Thr Ile Thr Val Lys 65 70 75 80

Glu Phe Leu Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly Glu Thr 85 90 95

Leu Ser His Ser His Leu Leu His Lys Lys Glu Asn Gly Ile Trp
100 105 110

Ser Thr Glu Ile Leu Lys Asn Phe Lys Asn Lys Thr Phe Leu Lys Cys 115 120 125

Glu Ala Pro Asn Tyr Ser Gly Arg Phe Thr Cys Ser Trp Leu Val Gln 130 135 140

Arg Asn Met Asp Leu Lys Phe Asn Ile Lys Ser Ser Ser Ser Pro 145 150 155 160

Asp Ser Arg Ala Val Thr Cys Gly Met Ala Ser Leu Ser Ala Glu Lys 165 170 175

Val Thr Leu Asp Gln Arg Asp Tyr Glu Lys Tyr Ser Val Ser Cys Gln 180 185 190

Glu Asp Val Thr Cys Pro Thr Ala Glu Glu Thr Leu Pro Ile Glu Leu 195 200 205

Ala Leu Glu Ala Arg Gln Gln Asn Lys Tyr Glu Asn Tyr Ser Thr Ser 210 215 220

Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro Lys Asn Leu Gln 225 230 235 240

Met Lys Pro Leu Lys Asn Ser Gln Val Glu Val Ser Trp Glu Tyr Pro 245 250 255 Asp Ser Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Lys Phe Phe Val 260 265 270

Arg Ile Gln Arg Lys Lys Glu Lys Met Lys Glu Thr Glu Glu Gly Cys 275 280 285

Asn Gln Lys Gly Ala Phe Leu Val Glu Lys Thr Ser Thr Glu Val Gln 290 295 300

Cys Lys Gly Gly Asn Val Cys Val Gln Ala Gln Asp Arg Tyr Tyr Asn 305 310 315 320

Ser Ser Cys Ser Lys Trp Ala Cys Val Pro Cys Arg Val Arg Ser 325 330 335

<210> 10

<211> 531

<212> PRT

<213> Mus musculus

<400> 10

Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp 1 5 10 15

Tyr Lys Asp Asp Asp Lys Leu Met Trp Glu Leu Glu Lys Asp Val 20 25 30

Tyr Val Val Glu Val Asp Trp Thr Pro Asp Ala Pro Gly Glu Thr Val 35 40 45

Asn Leu Thr Cys Asp Thr Pro Glu Glu Asp Asp Ile Thr Trp Thr Ser 50 55 60

Asp Gln Arg His Gly Val Ile Gly Ser Gly Lys Thr Leu Thr Ile Thr 65 70 75 80

Val Lys Glu Phe Leu Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly 85 90 95

Glu Thr Leu Ser His Ser His Leu Leu Leu His Lys Lys Glu Asn Gly
100 105 110

Ile Trp Ser Thr Glu Ile Leu Lys Asn Phe Lys Asn Lys Thr Phe Leu 115 120 125

Lys Cys Glu Ala Pro Asn Tyr Ser Gly Arg Phe Thr Cys Ser Trp Leu 130 135 140 5

Val Gln Arg Asn Met Asp Leu Lys Phe Asn Ile Lys Ser Ser Ser Ser Ser Pro Asp Ser Arg Ala Val Thr Cys Gly Met Ala Ser Leu Ser Ala Glu Lys Val Thr Leu Asp Gln Arg Asp Tyr Glu Lys Tyr Ser Val Ser Cys Gln Glu Asp Val Thr Cys Pro Thr Ala Glu Glu Thr Leu Pro Ile Glu Leu Ala Leu Glu Ala Arg Gln Gln Asn Lys Tyr Glu Asn Tyr Ser 210 Thr Ser Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro Lys Asn Leu Gln Met Lys Pro Leu Lys Asn Ser Gln Val Glu Val Ser Trp Glu 245 Tyr Pro Asp Ser Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Lys Phe Phe Val Arg Ile Gln Arg Lys Lys Glu Lys Met Lys Glu Thr Glu Glu 280 275 Gly Cys Asn Gln Lys Gly Ala Phe Leu Val Glu Lys Thr Ser Thr Glu Val Gln Cys Lys Gly Gly Asn Val Cys Val Gln Ala Gln Asp Arg Tyr Tyr Asn Ser Ser Cys Ser Lys Trp Ala Cys Val Pro Cys Arg Val Arg 330 Ser Ser Arg Gly Gly Ser Gly Ser Gly Gly Gly Gly Ser 340 345 350 Lys Leu Leu Ala Val Pro Arg Ser Ser Pro Asp Trp Ala Gln Cys Gln Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His Ala 370 375 380

Pro Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu Thr

Lys Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro Gln 405 410 415

Gly Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln Gly
420 425 430

Leu Val Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly Glu
435 440 445

Pro Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser Leu 450 455 460

Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu Thr 465 470 475 480

Gln Gln Met Pro Ser Leu Ser Ser Gln Gln Trp Gln Arg Pro Leu 485 490 495

Leu Arg Ser Lys Ile Leu Arg Ser Leu Gln Ala Phe Leu Ala Ile Ala 500 505 510

Ala Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu Val 515 520 525

Pro Thr Ala 530

<210> 11

<211> 521 <212> PRT

<213> Homo sapiens

<400> 11

Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp 1 5 10 15

Tyr Lys Asp Asp Asp Lys Leu Ile Trp Glu Leu Lys Lys Asp Val 20 25 30

Tyr Val Val Glu Leu Asp Trp Tyr Pro Asp Ala Pro Gly Glu Met Val 35 40 45

Val Leu Thr Cys Asp Thr Pro Glu Glu Asp Gly Ile Thr Trp Thr Leu 50 55 60

Asp Gln Ser Ser Glu Val Leu Gly Ser Gly Lys Thr Leu Thr Ile Gln 65 70 75 80

Val Lys Glu Phe Gly Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly 85 90 95

Glu Val Leu Ser His Ser Leu Leu Leu Leu His Lys Lys Glu Asp Gly
100 105 110

Ile Trp Ser Thr Asp Ile Leu Lys Asp Gln Lys Glu Pro Lys Asn Lys 115 120 125

Thr Phe Leu Arg Cys Glu Ala Lys Asn Tyr Ser Gly Arg Phe Thr Cys 130 135 140

Trp Trp Leu Thr Thr Ile Ser Thr Asp Leu Thr Phe Ser Val Lys Ser 145 150 155 160

Ser Arg Gly Ser Ser Asp Pro Gln Gly Val Thr Cys Gly Ala Ala Thr 165 170 175

Leu Ser Ala Glu Arg Val Arg Gly Asp Asn Lys Glu Tyr Glu Tyr Ser 180 185 190

Val Glu Cys Gln Glu Asp Ser Ala Cys Pro Ala Ala Glu Glu Ser Leu 195 200 205

Pro Ile Glu Val Met Val Asp Ala Val His Lys Leu Lys Tyr Glu Asn 210 215 220

Tyr Thr Ser Ser Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro 225 230 235 240

Asn Asn Leu Gln Leu Lys Pro Leu Lys Asn Ser Arg Gln Val Glu Val 245 250 255

Ser Trp Glu Tyr Pro Asp Thr Trp Ser Thr Pro His Ser Tyr Phe Ser 260 265 270

Leu Thr Phe Cys Val Gln Val Gln Gly Lys Ser Lys Arg Glu Lys Lys 275 280 285

Asp Arg Val Phe Thr Asp Lys Thr Ser Ala Thr Val Ile Cys Arg Lys 290 295 300

Asn Ala Ser Ile Ser Val Arg Ala Gln Asp Arg Tyr Tyr Ser Ser Ser 305 310 315 320

Trp Ser Glu Trp Ala Ser Val Pro Cys Ser Gly Ser Gly Ser Ser Arg 325 330 335

Gly Gly Ser Gly Gly Gly Gly Gly Gly Ser Lys Leu Arg 340 345 350

Ala Val Pro Gly Gly Ser Ser Pro Ala Trp Thr Gln Cys Gln Gln Leu 355 360 365

Ser Gln Lys Leu Cys Thr Leu Ala Trp Ser Ala His Pro Leu Val Gly 370 375 380

His Met Asp Leu Arg Glu Glu Gly Asp Glu Glu Thr Thr Asn Asp Val 385 390 395 400

Pro His Ile Gln Cys Gly Asp Gly Cys Asp Pro Gln Gly Leu Arg Asp 405 . 410 415

Asn Ser Gln Phe Cys Leu Gln Arg Ile His Gln Gly Leu Ile Phe Tyr 420 425 430

Glu Lys Leu Leu Gly Ser Asp Ile Phe Thr Gly Glu Pro Ser Leu Leu 435 440 445

Pro Asp Ser Pro Val Ala Gln Leu His Ala Ser Leu Leu Gly Leu Ser 450 455 460

Gln Leu Leu Gln Pro Glu Gly His His Trp Glu Thr Gln Gln Ile Pro 465 470 475 480

Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu Leu Arg Phe Lys 485 490 495

Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala Ala Arg Val Phe 500 505 510

Ala His Gly Ala Ala Thr Leu Ser Pro 515 520